Figure 1A

MEPETSHVADSYEISSTSFVATDNTDSSIVYLAAEQVLTGP	***
1 50 ID NO:17 ME 50 ID NO:17 ME 50 ID NO:10 ME 50 ID NO:16 ME 50 ID NO:16 ME 50 ID NO:16 ME 50 ID NO:16 ME 50 ID NO:17 ME 50 ID NO:17 ME 50 ID NO:10	i i

Figure 1B

118 VLSARSSEFKSALAAAKKEKDSNUTAAV	180				. 6		0		à	i			240	- →	1		Z (י מ	מ
ARSSFEKSALAAAKKEKDSNNTAAV	,,	KDYEVO	! !) V E E V E V	1 :) / ¼	1 1 1 1	 	1 · 1 i 1 t	*			KIPELI	 	E	VAELT	KITELI	KITELLI
ARSSFEKSALAAAKKEKDSNNTAAV		I I		1 1	LlGGGG		LVPGG-	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	1 1 1 1 1	1				LAFIF	1		AASTE(AAWAE	AAMAE
ARSSFEKSALAAAKKEKDSNNTAAV		YLELEKE YLELEKE	!!!!!!!!!	1	RIBLRE	!	RIKMEE	1 1 1	!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!	1 1 1 1 1 1 1 1 1	*			MLEVLY	 		MAQVLE	NVEQMA	
ARSSFEKSALAAAKKEKDSNNTAAV ARSTEFYNLFAARAAAAGGGGEDGGE ARSTEFYNLFAARGRGGDGAGGGGGGGGGGGGG ARSTEFYNLFAARGRGGDGAAGGGGGGGGGGGGGG ARSTEFYNLFAARGRGGDGAAGGGGGGGGGGGGG ARSTEFYNLFAARGRGGDGAAGGGGGGGGGGGGGGGGGGGGGGGGGGGG		1 1 1	1 1 1 1 1 1 1			i 	TGGRE		1					PAVDE	1 1 1 1 1 1		HPAVAF	FPAIRF	0021PF
ARSSFEKSALAAAKKEKDSNUTAAV			1 1 1 1	1 1 1 1 1 1 1 1 1 1	 	 	GGGGEF	1 1 1 1 1	1 1 1	1 1 1				CHVACI	 - - - -		AHVGCI	PHDSC	
ARSSFEKSALAAAKKEKDSNI ARSSFELPGVFARRAAAAGGGGI ARSTFEYNLFAARGRGGDGAAGG			1	! 	EDGGE-	1 1 1 1 1 1	366666	1 1 1 1 1 1	1 1 1 1 1 1 1	1				CADENC	1 1 1	 	CVDEDC	CADPMC	CMGG AC
ARSSFFKSALAAAKK ARSTEFYNLFAARRAAAA ARSTEFYNLFAARRAAAA		EKDSN	1	1 1 1	AGGGGI	 - - - -	:DGAAG	1	1 1 1 1	1				PKGVSE	1 1	1	PKAACL	SANGG	0111100
ARSSEFKSALA ARSTEFKILPGVFA ARSTEFYILFA AR		AAKE		 	FEAAAA	1 1 1 1 1 1	ARGEGG	1 1 1 1 1 1 1	1 1 1 1 1	1 1 1 1				RVRPPI		1	RVGDL	KLRPA	ו גרת ת דיווי
ARSSEE APPEL APPEL ARSTEE ARST		KSKLA	1	1	PCVFA	!	TMLFA	!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!	:	!	*			AYVYSS	!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!	 	OYLYSG	SYLYIG	CELT TITE
		ARSSEE	 	!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!	INGUE	!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!	RSTE	 			* * *			VVTVLA	!!!!	1 1	LRLVLI	FLSLL(FEC
	121	10:17	10:2	8 :01	10:10	10:4	10:16	NO:12	40: 6	NO:14			181	NO:17	NO: 2	8 :ON	NO:10	NO: 4	
7 2 8 6 4 9 2 9 4 7 7 8 9 4 7														Q T	QH		ū	ID	
ID NO: 17 ID NO: 2 ID NO: 8 ID NO: 10 ID NO: 16 ID NO: 16 ID NO: 17		SEQ	SEQ	SEQ	SEQ	SEQ	SEQ	SEQ	SEÇ	SEQ				SEQ	SEQ	SEQ	SEQ	SEQ	

-DAELSLLGYLYIGKLRPAPDDVVSCADPMCPHDSCPPAIRFNVEQMYAAWAFKITELIS

SEQ ID NO:12 SEQ ID NO: 6 SEQ ID NO:14

Figure 1C

241 7 - £1Q9-HLLLVVD&ZZVIEDTLVIL 2	m 👄 :	4 DEQRELENFUDNTEVEDVERILGVA	301 17 LVKELIDKRKELGLEVPKVKKHVSNVHKALDSDDIELVKLLLKEDHTNLDDAC 2VRRMRRALDAADIELVKLMVMGEGLDLDAAL 8PVHEKRVRRIHRALDSDDVELVKLLLNESDITLDDAN 10 VIKQIIDARLSLGLISPENKGFPNKHVRRIHRALDSDDVELVRMLLTEGQTNLDDAF 4
10		SEQ ID NO: SEQ ID NO:1 SEQ ID NO:1	SEQ ID NO: 10 SE

* * * *

* * *

Figure 1D

SEQ ID NO: 17 ALHEAVATCHVETATOLLKLDLADVNHRNPRGYFVLHVAAMFKEPQLILSLLEKGASASE SEQ ID NO: 2 AVHTAVQHGRD7VXALLELGAADVN

SEÇ ID NO:17 ATLEGETALMIAKQATMAVECNNIPEQCKHSLKGRLCVEILEQEDKREQIPRDVPPSFAV LTADGQSAMSICRLTRMKDYNTKMEQGQESNKDRLCIDILDREMIRKPMAVEDSVTSPL VTFDGRKAVQISKRLTKQGDYFGVTEEGKPSPYDRLCIEILEQAERRDPQLGEASVSLAM LTADGQSAMSICRRITRMKDYNTKMEQGQESNKDRLCIDILDREMIRKPMAVEDSVTSPL ----EQGQESNKDRMCIDILEREMMRNFMTAEDSVTSPL LTADGESAIGICRRLTRAKDYNTKMEQGQESNYDRLCIDILEREMMRNPMAVEDAVTSPL * * * * 421 SEQ 1D NO: 8 1D NO:16 ID NO:10 ID NO:12 ID NO: 6 ID NO:14 SEQ ID NO: 2 ID NO: 4 SEQ SEQ SEQ SEQ

Figure 1E

AACELENTLLDLENEVALAQRLFFTEAQAAMEIAEMEGTCEFIVTSLEFDRLTGTKRTSP LADDLHMKLLYLENRVAFARLFFPAEAKVAMQIAQADTTEEFG-GIVAVAASTSGKLREV AGESLEGRILYLENRVALARIMFPMEAFVAMDIAQVDGTLEFNIGS-GANFPPERQRTTV LAPDLHMKLLYLENPVAFARLFFPAEAKVAMQIAQADTTPEFGIVFAASTSGKLKEV LAPDLHMKLLYLENPVAFARLFFPAEAKVAMQIAQADTTPEFGIVFAASTSGKLKEV	LADDLHMKLSTLENRVAFARLFFPAE	
2	6 14	
	SEÇ ID SEÇ ID	

542	GVKIAFFRILEEHQSRLKALSKTVELGKRFFPFCSAVLDQIMNCEFLTQLACGEDDT	1	DINETPYTONKFLESRVDALMKTVELGRRYFPHCSQVLDKFLEDDLFEGLDQFYLQRGTA	DINESPFIMKEEHLARMTALSKTVELGKRFFPPCSNVLDKIMDDETDPVSLGRDTS		DENETPYTONKFLRSRVDALMKTVELGRRYFPNCSQVLDKFLEUDLPDS PDALDLQNGTS	DLNETPVTQNKFLRSRVDALMKTVELGRRYFPMCSQVLDKFLEDDLPDSPDALDLQNGTS		DENETPVTKNKFERSRVDALAKTVELGRRYFPNCSQVLDKFLEDGLPDGLDAFQQQSGTP
5	NO:17	No: 2	NO: 8	NO:10	NO: 4	NO:16	NO:12	NO: 6	NO:14
				ID 1				ID	
	SEQ	SEQ	SEQ	SEQ	SEQ	SEQ	SEQ	SEQ	SEQ

Figure 1F

649	SEQ ID NO:17 ABFFLYKKØRYMELQETLFKAFSEDNLELGNSSLTDSTSSTSFSTGGKPSNRKLSHRPR	88	DEGEV-KPMPFCELKEDVLKAFSKDKAEGSVFSGLSSSSCSPPQKYAQR	AEKEKRFEDLQDVLQKAFHEDKEENDRSGLSSSSSSSSIGAIRPRR		DEÇNV-KRMRFCELKEDVRKAFSKERADNSMFSILSSSSSSPPPKVAKK	DEGMV-RRMPFCELKEDVRKAFSFDRADNSMFSILSSSSSSS		DEGOV-REMPRETEVEDVPKAYSHDTADNSMFSALSSNSSS	
601	AEFFLYFE			AEKE		DEÇMV-KRI	DEQNV-KRI		DEÇÇV-KM	
)9	NO:17	SEQ ID NO: 2	SEÇ ID NO: 8	SEQ ID NO:10	SEQ ID NO: 4	SEQ ID NO:16	SEQ ID NO:12	SEQ ID NO: 6	SEQ 1D NO:14	
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